

Sequence listing

<110> Epigenomics AG
<120> Method for investigating cytosine methylation in DNA
5 sequences by means of triplex-forming oligomers
<160> 3

<210> 1
<211> 143
10 <212> DNA
<213> HOMO SAPIENS

<400> 1
cggaggaaga aagaggaggg gctggctggt caccagaggg tggggcggac cgcgtgcgct 60
15 cggcggctgc ggagaggggg agagcaggca gcgggcggcg gggagcagca tggagccggc 60
ggcggggagc agcatggagc ctt 23

<210> 2
<211> 131
20 <212> DNA
<213> HOMO SAPIENS

<400> 2
cggaggaaga aagaggaggg gctggctggt caccagaggg tggggcggac cgcgtgcgct 60
25 cggcggctgc ggagaggggg agagcaggca gcgggcggcg gggagcagca tggagccggc 60
ggcggggagc a 11

<210> 3
<211> 143
30 <212> DNA
<213> HOMO SAPIENS

<400> 3
cggaggaaga aagaggaggg gctggctggt caccagaggg tggggcggac cgcgtgcgct 60
35 cggcggctgc ggagaggggg agagcaggca gcgggcggcg gggagcagca tggagccggc 60
ggcggggagc agcatggagc ctt 23